

## SEQUENCE LISTING

<110> Reactive Surfaces  
McDaniel, C. Steven

<120> Recombinant Organophosphorus Acid Anhydrase and Methods of Use

<130> RACT-00200

<140> Unknown

<141> 2002-12-28

<150> 07/928,540

<151> 1992-08-13

<150> 08/252,384

<151> 1994-06-01

<150> 07/344,258

<151> 1989-04-27

<160> 1

<170> PatentIn version 3.2

<210> 1

<211> 337

<212> PRT

<213> Pseudomonas aeruginosa

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Thr Ile Ser Glu Ala Gly Phe Thr Leu Thr His Glu His Ile Cys Gly  
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Ser Ser Ala Gly Phe Leu Arg Ala Trp Pro Glu Phe Phe Gly Ser Arg  
35 40 45

Lys Ala Leu Ala Glu Lys Ala Val Arg Gly Leu Arg Arg Ala Arg Ala  
50 55 60

Ala Gly Val Arg Thr Ile Val Asp Val Ser Thr Phe Asp Ile Gly Arg  
65 70 75 80

Asp Val Ser Leu Leu Ala Glu Val Ser Arg Ala Ala Asp Val His Ile  
85 90 95

Val Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu  
100 105 110

Arg Ser Val Glu Glu Leu Thr Gln Phe Phe Leu Arg Glu Ile Gln Tyr  
115 120 125

Gly Ile Glu Asp Thr Gly Ile Arg Ala Gly Ile Ile Lys Val Ala Thr  
130 135 140

Thr Gly Lys Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala  
145 150 155 160

Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala  
165 170 175

Ser Gln Arg Asp Gly Glu Gln Gln Ala Ala Ile Phe Glu Ser Glu Gly  
180 185 190

Leu Ser Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp Asp  
195 200 205

Leu Ser Tyr Leu Thr Ala Leu Ala Ala Arg Gly Tyr Leu Ile Gly Leu  
210 215 220

Asp His Ile Pro His Ser Ala Ile Gly Leu Glu Asp Asn Ala Ser Ala  
225 230 235 240

Ser Ala Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile  
245 250 255

*E3*  
*Conclude*

Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln Ile Leu Val Ser Asn  
260 265 270

Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val  
275 280 285


Met Asp Arg Val Asn Pro Asp Gly Met Ala Phe Ile Pro Leu Arg Val  
290 295 300

Ile Pro Phe Leu Arg Glu Lys Gly Val Pro Gln Glu Thr Leu Ala Gly  
305 310 315 320

Ile Thr Val Thr Asn Pro Ala Arg Phe Leu Ser Pro Thr Leu Arg Ala  
325 330 335

Ser

## SEQUENCE LISTING

 <110> Reactive Surfaces, Ltd.

<120> Recombinant Organophosphorus Acid Anhydrase and Methods of Use

<130> RACT-00200

<140> Unknown

<141> 2003-01-02

<150> 07/928,540

<151> 1992-08-13

<150> 08/252,384

<151> 1994-06-01

<150> 07/344,258

<151> 1989-04-27

<160> 2

<170> PatentIn version 3.1

<210> 1

<211> 1014

<212> DNA

<213> Pseudomonas diminuta

<220>

<221> CDS

<222> (1)..(1011)

<223>

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1 5 10 15

aca atc tct gaa gcg ggt ttc aca ctg act cac gag cac atc tgc ggc  
96  
Thr Ile Ser Glu Ala Gly Phe Thr Leu Thr His Glu His Ile Cys Gly

20

25

30

agc tcg gca gga ttc ttg cgt gct tgg cca gag ttc ttc ggt agc cgc  
144

Ser Ser Ala Gly Phe Leu Arg Ala Trp Pro Glu Phe Phe Gly Ser Arg

35

40

45

aaa gct cta gcg gaa aag gct gtg aga gga ttg cgc cgc gcc aga gcg  
192

Lys Ala Leu Ala Glu Lys Ala Val Arg Gly Leu Arg Arg Ala Arg Ala

50

55

60

gct ggc gtg cga acg att gtc gat gtg tcg act ttc gat atc ggt cgc  
240

Ala Gly Val Arg Thr Ile Val Asp Val Ser Thr Phe Asp Ile Gly Arg

65

70

75

80

gac gtc agt tta ttg gcc gag gtt tcg cgg gct gcc gac gtt cat atc  
288

Asp Val Ser Leu Leu Ala Glu Val Ser Arg Ala Ala Asp Val His Ile

85

90

95

gtg gcg gcg acc ggc ttg tgg ttc gac ccg cca ctt tcg atg cga ttg  
336

Val Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu

100

105

110

agg agt gta gag gaa ctc aca cag ttc ttc ctg cgt gag att caa tat  
384

Arg Ser Val Glu Glu Leu Thr Gln Phe Phe Leu Arg Glu Ile Gln Tyr

115

120

125

ggc atc gaa gac acc gga att agg gcg ggc att atc aag gtc gcg acc  
432

Gly Ile Glu Asp Thr Gly Ile Arg Ala Gly Ile Ile Lys Val Ala Thr

130

135

140

aca ggc aag gcg acc ccc ttt cag gag tta gtg tta aag gcg gcc gcc  
480

Thr Gly Lys Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala

145

150

155

160

cgg gcc agc ttg gcc acc ggt gtt ccg gta acc act cac acg gca gca  
528

Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala

165

170

175

agt cag cgc gat ggt gag cag cag gcc gcc att ttt gag tcc gaa ggc  
576

Ser Gln Arg Asp Gly Glu Gln Gln Ala Ala Ile Phe Glu Ser Glu Gly

180

185

190

ttg agc ccc tca cgg gtt tgt att ggt cac agc gat gat act gac gat  
624

Leu Ser Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp Asp

195

200

205

ttg agc tat ctc acc gcc ctc gct gcg cgc gga tac ctc atc ggt cta  
672

Leu Ser Tyr Leu Thr Ala Leu Ala Ala Arg Gly Tyr Leu Ile Gly Leu

210

215

220

gac cac atc ccg cac agt gcg att ggt cta gaa gat aat gcg agt gca  
720

Asp His Ile Pro His Ser Ala Ile Gly Leu Glu Asp Asn Ala Ser Ala

225

230

235

240

tca gcc ctc ctg gcc atc cgt tcg tgg caa aca cgg gct ctc ttg atc  
768

Ser Ala Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile  
245 250 255

aag gcg ctc atc gac caa ggc tac atg aaa caa atc ctc gtt tgc aat  
816  
Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln Ile Leu Val Ser Asn  
260 265 270

gac tgg ctg ttc ggg ttt tgc agc tat gtc acc aac atc atg gac gtg  
864  
Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val  
275 280 285

atg gat cgc gtg aac ccc gac ggg atg gcc ttc att cca ctg aga gtg  
912  
Met Asp Arg Val Asn Pro Asp Gly Met Ala Phe Ile Pro Leu Arg Val  
290 295 300

atc cca ttc gta cga gag aag ggc gtc cca cag gaa acg ctg gca ggc  
960  
Ile Pro Phe Val Arg Glu Lys Gly Val Pro Gln Glu Thr Leu Ala Gly  
305 310 315 320

atc act gtg act aac ccg gcg cgg ttc tat gtc acc gac ctt gcg ggc  
1008  
Ile Thr Val Thr Asn Pro Ala Arg Phe Tyr Val Thr Asp Leu Ala Gly  
325 330 335

gtc atg  
1014  
Val

<210> 2  
<211> 337  
<212> PRT  
<213> Pseudomonas diminuta

<400> 2

Met Ser Ile Gly Thr Gly Asp Arg Ile Asn Thr Val Arg Gly Pro Ile  
1 5 10 15

Thr Ile Ser Glu Ala Gly Phe Thr Leu Thr His Glu His Ile Cys Gly  
20 25 30

Ser Ser Ala Gly Phe Leu Arg Ala Trp Pro Glu Phe Phe Gly Ser Arg  
35 40 45

Lys Ala Leu Ala Glu Lys Ala Val Arg Gly Leu Arg Arg Ala Arg Ala  
50 55 60

Ala Gly Val Arg Thr Ile Val Asp Val Ser Thr Phe Asp Ile Gly Arg  
65 70 75 80

Asp Val Ser Leu Leu Ala Glu Val Ser Arg Ala Ala Asp Val His Ile  
85 90 95

Val Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu  
100 105 110

Arg Ser Val Glu Glu Leu Thr Gln Phe Phe Leu Arg Glu Ile Gln Tyr  
115 120 125

Gly Ile Glu Asp Thr Gly Ile Arg Ala Gly Ile Ile Lys Val Ala Thr  
130 135 140

Thr Gly Lys Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala  
145 150 155 160

Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala



165

170

175

Ser Gln Arg Asp Gly Glu Gln Gln Ala Ala Ile Phe Glu Ser Glu Gly  
180 185 190

Leu Ser Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp Asp  
195 200 205

Leu Ser Tyr Leu Thr Ala Leu Ala Ala Arg Gly Tyr Leu Ile Gly Leu  
210 215 220

Asp His Ile Pro His Ser Ala Ile Gly Leu Glu Asp Asn Ala Ser Ala  
225 230 235 240

Ser Ala Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile  
245 250 255

Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln Ile Leu Val Ser Asn  
260 265 270

Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val  
275 280 285

Met Asp Arg Val Asn Pro Asp Gly Met Ala Phe Ile Pro Leu Arg Val  
290 295 300

Ile Pro Phe Val Arg Glu Lys Gly Val Pro Gln Glu Thr Leu Ala Gly  
305 310 315 320

Ile Thr Val Thr Asn Pro Ala Arg Phe Tyr Val Thr Asp Leu Ala Gly  
325 330 335

Val

Protein Comparisons  
Percent Identity

McDaniel vs Wild Lab 57.8%  
McDaniel vs Serdar 59.5%  
Serdar vs Wild Lab 100%

DNA Comparisons  
Percent Identity

McDaniel vs Wild Lab 88.0%  
McDaniel vs Serdar 83.7%  
Serdar vs Wild Lab 99.6%

McDaniel.pro Met Gln Thr Arg Arg Val Val Leu Lys Ser Ala Ala Ala Arg Thr Leu Leu Gly Gly Leu Ala Gly Cys Ala Thr Trp Leu Asp Arg 29  
Wild Lab.PRO Met Gln Thr Arg Arg Val Val Leu Lys Ser Ala Ala Ala Arg Thr Leu Leu Gly Gly Leu Ala Gly Cys Ala Thr Trp Leu Asp Arg 29  
Serdar.pro Met Gln Thr Arg Arg Val Val Leu Lys Ser Ala Ala Ala Arg Thr Leu Leu Gly Gly Leu Ala Gly Cys Ala Thr Trp Leu Asp Arg 30

McDaniel.pro Ser Ala Gln Ala Met Arg Ser Ile Arg Ala Arg Pro Ile Thr Ile Ser Glu Ala Gly Phe Thr Leu Thr His Glu Asp Ile Ser Ala 58  
Wild Lab.PRO Ser Ala Gln Ala Met Arg Ser Ile Arg Ala Arg Pro Ile Thr Ile Ser Glu Ala Gly Phe Thr Leu Thr His Glu Asp Ile Ser Ala 32  
Serdar.pro Ser Ala Gln Ala Met Arg Ser Ile Arg Ala Arg Pro Ile Thr Ile Ser Glu Ala Gly Phe Thr Leu Thr His Glu Asp Ile Ser Ala 60

McDaniel.pro Ala Arg Gln Asp Ser Cys Val Leu Gly Gln Ser Ser Ser Val Ala Cln Ser Ser Ser Gly Lys Gly Cys Glu Arg 83  
Wild Lab.PRO Ala Arg Gln Asp Ser Cys Val Leu Gly Gln Ser Ser Ser Val Ala Cln Ser Ser Ser Gly Lys Gly Cys Glu Arg 60  
Serdar.pro Ala Arg Gln Asp Ser Cys Val Leu Gly Gln Ser Ser Ser Val Ala Cln Ser Ser Ser Gly Lys Gly Cys Glu Arg 88

McDaniel.pro Ile Ala Arg Gln Ser Gly Trp Arg Ala Asn Asp Cys Arg Cys Val Asp Phe Arg Tyr Arg Ser Arg Arg Gln Phe Ile Gly Arg Gly Phe 11  
Wild Lab.PRO Ile Ala Arg Gln Ser Gly Trp Arg Ala Asn Asp Cys Arg Cys Val Asp Phe Arg Tyr Arg Ser Arg Arg Gln Phe Ile Gly Arg Gly Phe 89  
Serdar.pro Ile Ala Arg Gln Ser Gly Trp Arg Ala Asn Asp Cys Arg Cys Val Asp Phe Arg Tyr Arg Ser Arg Arg Gln Phe Ile Gly Arg Gly Phe 11'

McDaniel.pro Ala Gly Cys Arg Arg Ser Tyr Leu Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu Arg Tyr Val Glu Glu Leu Thr 14  
Wild Lab.PRO Ala Gly Cys Arg Arg Ser Tyr Leu Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu Arg Tyr Val Glu Glu Leu Thr 11  
Serdar.pro Ala Gly Cys Arg Arg Ser Tyr Leu Ala Ala Thr Gly Leu Trp Phe Asp Pro Pro Leu Ser Met Arg Leu Arg Tyr Val Glu Glu Leu Thr 14'

McDaniel.pro Leu Val Leu Pro Ala Val Arg Phe Asn Met Ala Ser Lys Tyr Thr Gly Ile Arg Ala Gly Ile Ile Lys Val Ala Thr Thr Gly Lys 17  
Wild Lab.PRO Leu Val Leu Pro Ala Val Arg Phe Asn Met Ala Ser Lys Tyr Thr Gly Ile Arg Ala Gly Ile Ile Lys Val Ala Thr Thr Gly Lys 14'  
Serdar.pro Leu Val Leu Pro Ala Val Arg Phe Asn Met Ala Ser Lys Tyr Thr Gly Ile Arg Ala Gly Ile Ile Lys Val Ala Thr Thr Gly Lys 17

McDaniel.pro Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala Ser 20  
Wild Lab.PRO Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala Ser 17  
Serdar.pro Ala Thr Pro Phe Gln Glu Leu Val Leu Lys Ala Ala Ala Arg Ala Ser Leu Ala Thr Gly Val Pro Val Thr Thr His Thr Ala Ala Ser 20

McDaniel.pro Gln Arg Asp Gly Glu Arg Gly Arg Pro Pro Phe Leu Ser Pro Lys Leu Glu Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp 23  
Wild Lab.PRO Gln Arg Asp Gly Glu Arg Gly Arg Pro Pro Phe Leu Ser Pro Lys Leu Glu Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp 20  
Serdar.pro Gln Arg Asp Gly Glu Arg Gly Arg Pro Pro Phe Leu Ser Pro Lys Leu Glu Pro Ser Arg Val Cys Ile Gly His Ser Asp Asp Thr Asp 23

McDaniel.pro Asp Leu Ser Tyr Leu Thr Ala Leu Leu Arg Gly Tyr Leu Ile Gly Leu Asp His Ile Pro His Ser Ala Ile Gly Leu Glu Asp Asn 26  
Wild Lab.PRO Asp Leu Ser Tyr Leu Thr Ala Leu Leu Arg Gly Tyr Leu Ile Gly Leu Asp His Ile Pro His Ser Ala Ile Gly Leu Glu Asp Asn 23  
Serdar.pro Asp Leu Ser Tyr Leu Thr Ala Leu Leu Arg Gly Tyr Leu Ile Gly Leu Asp His Ile Pro His Ser Ala Ile Gly Leu Glu Asp Asn 26

McDaniel.pro Ala Ser Ala Ser Pro Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln 29  
Wild Lab.PRO Ala Ser Ala Ser Pro Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln 26  
Serdar.pro Ala Ser Ala Ser Pro Leu Leu Gly Ile Arg Ser Trp Gln Thr Arg Ala Leu Leu Ile Lys Ala Leu Ile Asp Gln Gly Tyr Met Lys Gln 29

McDaniel.pro Ile Leu Val Ser Asn Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val Met Asp Arg Val Asn Pro Asp Gly Met 32  
Wild Lab.PRO Ile Leu Val Ser Asn Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val Met Asp Arg Val Asn Pro Asp Gly Met 29  
Serdar.pro Ile Leu Val Ser Asn Asp Trp Leu Phe Gly Phe Ser Ser Tyr Val Thr Asn Ile Met Asp Val Met Asp Arg Val Asn Pro Asp Gly Met 32

McDaniel.pro Ala Phe Ile His 32  
Wild Lab.PRO Ala Phe Ile His 32  
Serdar.pro Ala Phe Ile His 35

McDaniel.pro 32  
Wild Lab.PRO Arg Phe Leu Ser Pro Thr Leu Arg Ala Ser 33  
Serdar.pro Arg Phe Leu Ser Pro Thr Leu Arg Ala Ser 36

Decoration 'Decoration #1': Shade (with black at 10% fill) residues that differ from McDaniel.pro.

Decoration 'Decoration #2': Box residues that differ from McDaniel.pro.

[illegible]

Decorations 'Decorations #1': Shaded (with black as 100 fill) residues that differ from McDaniel P diminuta.ssq.

Decorations 'Decorations 02': Box residues that differ from McDaniel P diminuta.ssq.

**Works Cited**

**McDaniel P. diminuta :**

McDaniel, C.S., Harper, L.L. and Wild, J.R., Cloning and sequencing of a plasmid-borne gene (opd) encoding a phosphotriesterase, J. Bacteriol. 170 (5), 2306-2311 (1988). Gene Bank Accession Number: M20392

**Oph-lab RC:** Wild lab DNA sequence

**Serdar:** Serdar Sequence obtained from United States Patent and Trademark Office (uspto.gov) Patent Number 5,484,728

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FILED 12-23-02 AND 01-02-03.